

Fig. 1a

PGIA-01-A8	EVQLLESGGGLVQPGGSLRLSCAASGFTFS	SYAMS	WVRQAPGKGLEWVS	A	50
Vh3_DP-47__3-23_	EVQLLESGGGLVQPGGSLRLSCAASGFTFS	SYAMS	WVRQAPGKGLEWVS	A	
PGIA-01-A8	ISGSGGSTYYADSVKG	RFTISRDN SKNTLYLQMN SLRAEDTAVYYCAK	DH	100	
Vh3_DP-47__3-23_	ISGSGGSTYYADSVKG	RFTISRDN SKNTLYLQMN SLRAEDTAVYYCAK	.		
PGIA-01-A8	NYDSSGYLDY	WGQGT LVT VSS	121	SEQ ID NO:140	
Vh3_DP-47__3-23_	WGQGT LVT VSS	JH1/JH4/JH5	SEQ ID NO:154	
PGIA-01-A8	NFMLTQPHSVSESPGKTVTISC	TRSSGSI AFDYVQ	WYQQRPGSAPT TVIY	50	
Vlambda6_6a	NFMLTQPHSVSESPGKTVTISC	TRSSGSI ASNYVQ	WYQQRPGSSPT TVIY		
PGIA-01-A8	EDNQRP S	GVPDRF S	ASIDSSSNSASLTIS	ALKTEDEADYYC	100
Vlambda6_6a	EDNQRP S	GVPDRF S	GSIDSSSNSASLTIS	GLKTEDEADYYC	
PGIA-01-A8	V	FGGGTKLTVL	111	SEQ ID NO:141	
Vlambda6_6a	:	FGGGTKLTVL	JL2/JL3	SEQ ID NO:158	

Fig. 1b

PGIA-03-A9	QVQLQESGPGLVKPSGTLSTLCAVSGGSIS	TSDWWS	WVRPPPGKGLEWIG	50
Vh4_DP-70__4-04_	QVQLQESGPGLVKPSGTLSTLCAVSGGSIS	SSNWS	WVRQPPGKGLEWIG	
PGIA-03-A9	EIYHSGSTNYHPSLKS	RVTISL	DKSKNQFSLKLSVTAADTAVYCAR	EG 100
Vh4_DP-70__4-04_	EIYHSGSTNYNP SLKS	RVTISV	DKSKNQFSLKLSVTAADTAVYCAR	..
PGIA-03-A9	GHSGSYPLIDY	WGKGT	LVTVSS	121
Vh4_DP-70__4-04_	WGQGT	LVTVSS	JH4
				SEQ ID NO:142
				SEQ ID NO:155
PGIA-03-A9	NFMLTQPHSVSESPGKTVTISC	TRSSGSIASNYVQ	WYQQRPGSSPTTVIY	50
Vlambda6_6a	NFMLTQPHSVSESPGKTVTISC	TRSSGSIASNYVQ	WYQQRPGSSPTTVIY	
PGIA-03-A9	EDNQRPS	GVPDRFSGSIDSSNSASLTISGLKTEDEADYYC	QSYDSSNOG	100
Vlambda6_6a	EDNQRPS	GVPDRFSGSIDSSNSASLTISGLKTEDEADYYC	QSYDSSN...	
PGIA-03-A9	VV	FGG	TKLTVL	112
Vlambda6_6a	..	FGG	TKLTVL	JL2/JL3
				SEQ ID NO:143
				SEQ ID NO:158

Fig 1c

PGIA-03-A11	QVQLVQSGG PE EVKKPGASV EV SKKASGYTFT	GDYMH	WVRQAPGQG PE EWMG	W	50
Vh1_DP-8_75__1-02__	QVQLVQSG AE EVKKPGASV KV SKKASGYTFT	GYVMH	WVRQAPGQG LE EWMG	W	
PGIA-03-A11	INPQTGVTKYAQKFQ	RVT M ARDT SIN TAYMEL RGL RSDDTAVY CV R	ED	I	100
Vh1_DP-8_75__1-02__	INPNSGGTNYAQKFQ	RVT M TRDT SIS TAYMEL SR LRSDDDTAVY CA R			
PGIA-03-A11	HNYDLMSAYNGLDV	WGQGTILVTVSS			125
Vh1_DP-8_75__1-02__	WGQGTILVTVSS			JH1
					SEQ ID NO:144
					SEQ ID NO:156
PGIA-03-A11	QSVLTQPPSVSAAPGQKV TISC	SGSSSNIGNNHVS	WYQQL AG TAPKLL IF		50
Vlambda1_DPL5__1b__	QSVLTQPPSVSAAPGQKV TISC	SGSSSNIGNNXVS	WYQQL PG TAPKLL IY		
PGIA-03-A11	DNDKRPS	GIPDRFSGSKSGTSATLGITGLQTGDEADY YC	GTWDKSPTDIY		100
Vlambda1_DPL5__1b__	DNNKRPS	GIPDRFSGSKSGTSATLGITGLQTGDEADY YC	GTWDSLSA		
PGIA-03-A11	V	FG S GTKLTVL			111
Vlambda1_DPL5__1b__	E	FG T GTKVTVL			JL1
					SEQ ID NO:145
					SEQ ID NO:159

Fig. 1d

PGIA-03-B2	QVQLQESGPGLVKPSA	TL	SLTCAVGG	SSIS	SNHWNS	WVRQSPGKGLEWIG	50
Vh4_DP-70__4-04_	QVQLQESGPGLVKPSG	TL	SLTCAVGG	SSIS	SSNWNNS	WVRQPPGKGLEWIG	
PGIA-03-B2	EIYTYGGAGNYNP	SLKS	RV	DISMDKSKNQFSL	HL	SSVTAADTAVYCGR	100
Vh4_DP-70__4-04_	EIYHSGSTNYNP	SLKS	RV	TISVDKSKNQFSL	KL	SSVTAADTAVYCAR	..
PGIA-03-B2	TGYDCFDI	WGQGT	LVTVSS	119		SEQ ID NO:148	
Vh4_DP-70__4-04_	WGQGT	LVTVSS	JH4		SEQ ID NO:155	
PGIA-03-B2	QAVLTQPPSSVSGAPGQ	RV	TISC	TGSSSNIGAGYDVH	WYQQLPGTAPKLLI	50	
Vlambda1_DPL8__1e_	QSVLTQPPSVSGAPGQ	RV	TISC	TGSSSNIGAGYDVH	WYQQLPGTAPKLLI		
PGIA-03-B2	Y GNSNRPS	GVPDRFSGSKSGTSASLA	ITGLQAEDEADYYC	QSYDSSLGV	100		
Vlambda1_DPL8__1e_	Y GNSNRPS	GVPDRFSGSKSGTSASLA	ITGLQAEDEADYYC	QSYDSSLG.			
PGIA-03-B2	FGTGTQLTVL	110			SEQ ID NO:149		
Vlambda1_DPL8__1e_	FGGGTQLTVL	JL7			SEQ ID NO:160		

Fig 1e

PGIA-04-A5	QVQLQESGPGLVKPSGTL SL TCAVSGGSIS	TSDWWS WVR R PPGKGLEWIG	50
Vh4_DP-70__4-04_	QVQLQESGPGLVKPSGTL SL TCAVSGGSIS	SSNWWS WVR Q PPGKGLEWIG	
PGIA-04-A5	EIYHSGSTNYHP SLKS RVTIS LD DKSKNQFSLKLSVTAADTAVYYCAR	EG 100	
Vh4_DP-70__4-04_	EIYHSGSTNYNP SLKS RVTIS VD DKSKNQFSLKLSVTAADTAVYYCAR	.	
PGIA-04-A5	GHSGSYPLDY WGRGTLVTVSS 121	SEQ ID NO:150	
Vh4_DP-70__4-04_ WGRGTLVTVSS JH2	SEQ ID NO:155	
PGIA-04-A5	NFMLTQPHSVSESPGKTA T IS	TGSGGSIARS YVQ WYQQRPG RAPS I V IY	50
Vlambda6_6a	NFMLTQPHSVSESPGKT V TIS	TRSSGSIASN YVQ WYQQRPG SSPT T V IY	
PGIA-04-A5	EDYQ RPS GVPDRFSGSIDSSSNSASLTITGLKT D EADYYC	QSSDDNNNV	100
Vlambda6_6a	EDNQ RPS GVPDRFSGSIDSSSNSASLTISGLKT E EADYYC	QSYDSSN ..	
PGIA-04-A5	V FGGGTKVTVL 111	SEQ ID NO:151	
Vlambda6_6a	V FGGGTKVTVL JL2/JL3	SEQ ID NO:158	

Fig 1f

PGIA-04-A8	QVQLQESGPGLVKPSSETLSLTCTNVSGGSIR	NYFWS	WIRQPPGQGLEWIG	Y	50
Vh4_DP-71__4-59_	QVQLQESGPGLVKPSSETLSLTCTVSGGSIS	SYFWS	WIRQPPGKGLEWIG	Y	
PGIA-04-A8	IYYSGTTDYNPSLKG	RVTISL	DTSKTQFSLKLSVTAADTA	FYYCVR	GPN 100
Vh4_DP-71__4-59_	IYYSGSTNYPNPSLKS	RVTISV	DTSKNQFSLKLSVTAADTA	VYYCAR	...
PGIA-04-A8	KYAFDP	WGQGLVTVSS	117		SEQ ID NO:152
Vh4_DP-71__4-59_	WGQGLVTVSS	JH4		SEQ ID NO:157
PGIA-04-A8	SYELTQPPSVSVSPGQTASITC	SGDKLGDKFAS	WYQQK	AGQSPVLVIY	RD 50
Vlambda3_DPL23__3r_	SYELTQPPSVSVSPGQTASITC	SGDKLGDKYAC	WYQQK	PGQSPVLVIY	QD
PGIA-04-A8	TKRPS	GIPERFSGNSGNTATLTISGTQAMDEADYYC	QAWDSSTAV	FGTG	100
Vlambda3_DPL23__3r_	SKRPS	GIPERFSGNSGNTATLTISGTQAMDEADYYC	QAWDSSTA.	FGTG	
PGIA-04-A8	TKVTVL	106			SEQ ID NO:153
Vlambda3_DPL23__3r_	TKVTVL	JL1			SEQ ID NO:161

Fig. 1g

PGIA-05-A1	QQLQESGPGLVKPSGTLTLTCAVSGGSIS	TSDWWS	WVRPPPGKLEWIG	50
Vh4_DP-70__4-04_	QVQLQESGPGLVKPSGTLTLTCAVSGGSIS	SSNWWS	WVRQPPGKLEWIG	
PGIA-05-A1	EIYHSGSTNYHPSLKS	RVTISL	DKSKNQFSLKLSVTAADTAVYCAR	EG 100
Vh4_DP-70__4-04_	EIYHSGSTNYNP	SLKS	RVTISV	DKSKNQFSLKLSVTAADTAVYCAR ..
PGIA-05-A1	GHSGSYPLIDY	WGRGTLVTVSS	121	SEQ ID NO:146
Vh4_DP-70__4-04_	WGRGTLVTVSS	JH2	SEQ ID NO:155
PGIA-05-A1	NFMLTQPHSVSESPGKTVTISC	ARSSGSIASNYVQ	WYQQRPGSSPTTIY	50
Vlambda6_6a	NFMLTQPHSVSESPGKTVTISC	TRSSGSIASNYVQ	WYQQRPGSSPTTVIY	
PGIA-05-A1	EDRQRPS	GVPDRFSGSIDSSSNSASLTISGLKTEDEADYYC	QSYDSSDHV	100
Vlambda6_6a	EDNQRPS	GVPDRFSGSIDSSSNSASLTISGLKTEDEADYYC	QSYDSSN	
PGIA-05-A1	V FGGGTKLTVL	111	SEQ ID NO:147	
Vlambda6_6a	V FGGGTKLTVL	JL2/JL3	SEQ ID NO:158	

Figure 2

c-Met IgG Antibodies In Europium Ligand Competition Assay

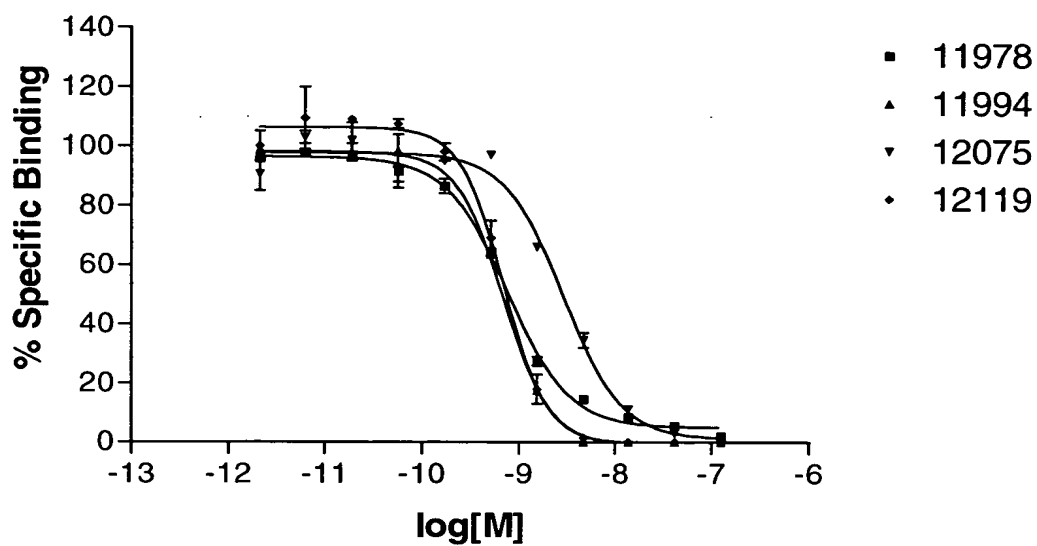


Figure 3

Inhibition of 184B5 Proliferation (0.133 nM rhHGF) by IgG

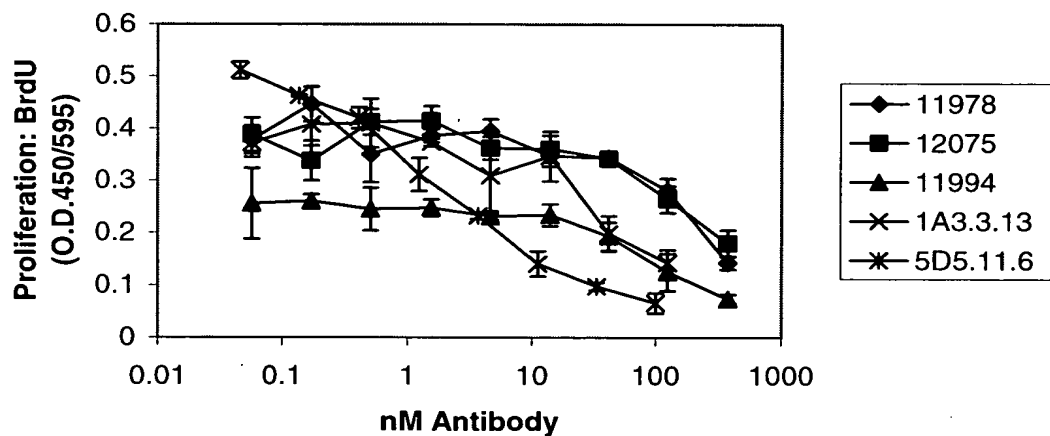


Figure 4

Tyrosine Phosphorylation of c-Met by c-Met IgG Antibodies

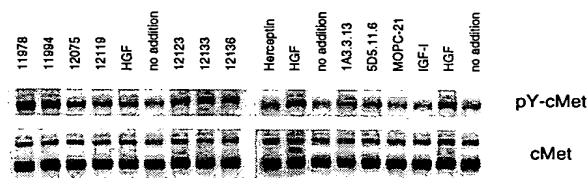
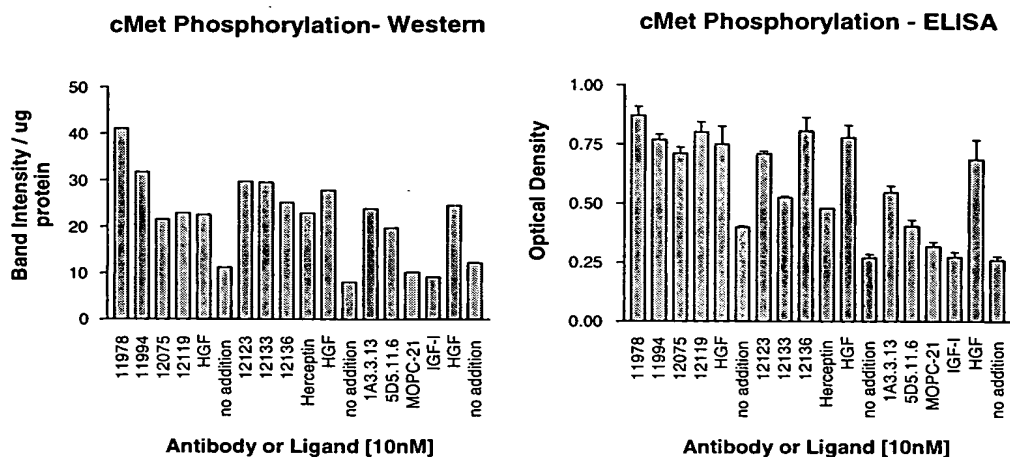


Figure 5

c-Met Fab in Europium Ligand Competition Assay

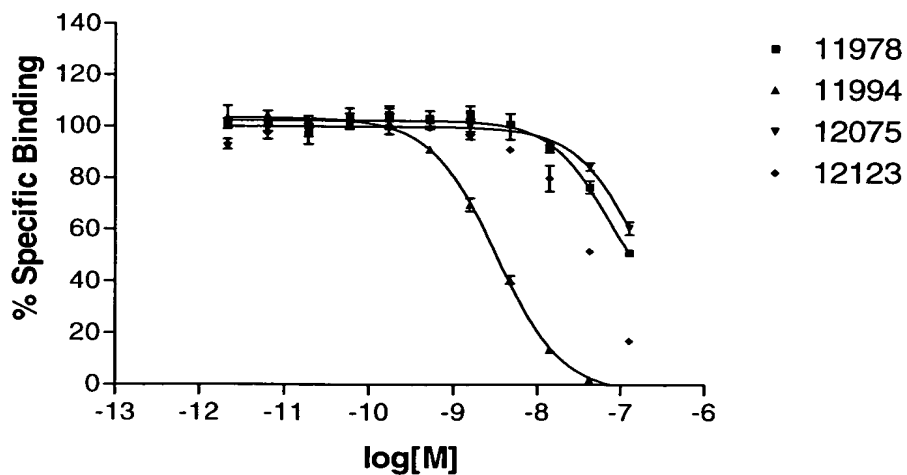


Figure 6

Tyrosine Phosphorylation of c-Met by c-Met Fab

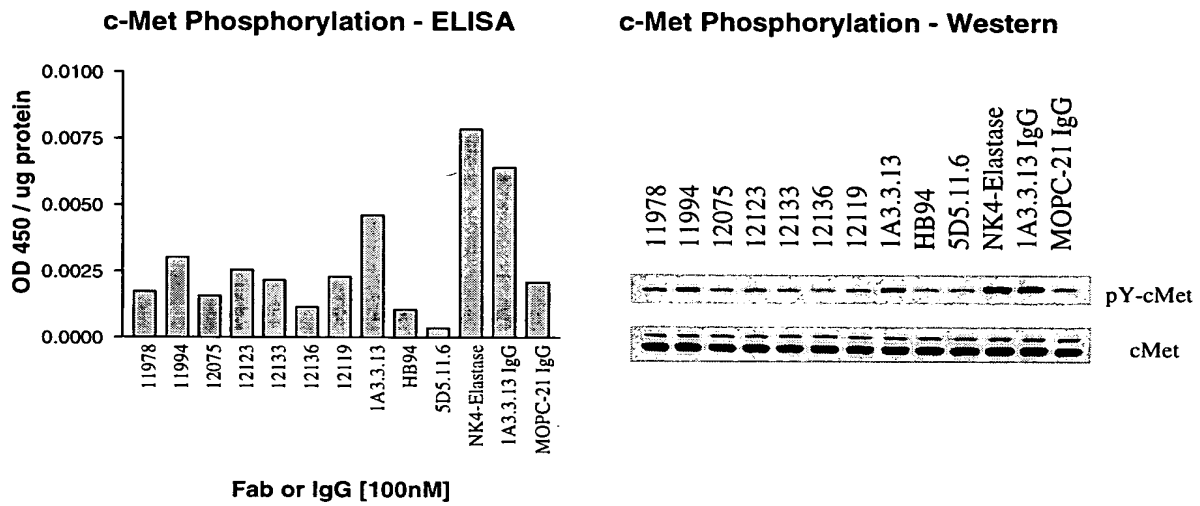


Figure 7

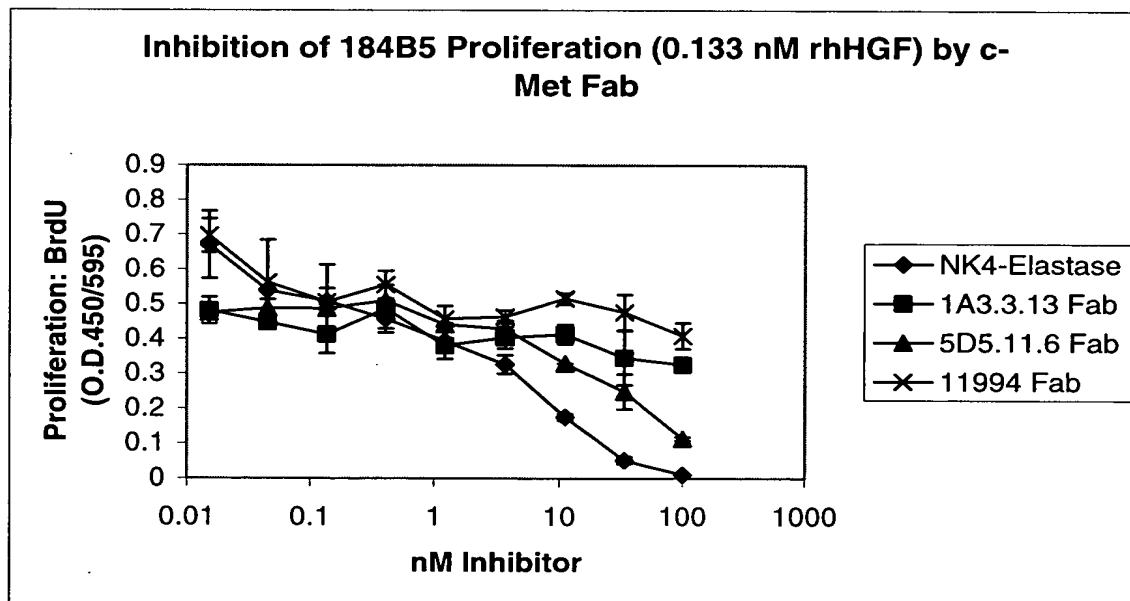


Figure 8

**DU145 Scatter Assay With A
c-Met IgG Antibody**

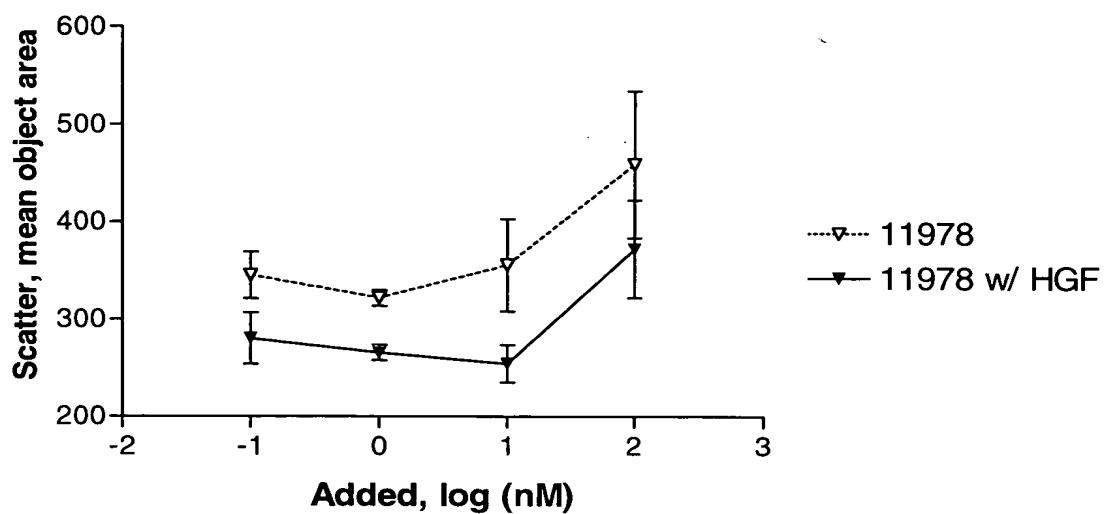


Fig. 9

**Inhibition of HGF Induced Migration by anti c-met Antibodies
in the Presence of 225pM HGF**

